Sequence Listing

<110> Botstein, David

Desnoyers,Luc

Ferrara, Napoleone

Fong, Sherman

Gao, Wei-Qiang

Goddard, Audrey

Gurney, Austin L.

Pan, James

Roy, Margaret Ann

Stewart, Timothy A.

Tumas, Daniel

Watanabe, Colin K.

Wood, William I.

<120> Secreted and Transmembrane Polypeptides and Nucleic Acids Encoding the Same

<130> P2930R1C10

<150>60/095,325

<151>1998-08-04

<150> 60/112,851

<151> 1998-12-16

<150> 60/113,145

<151> 1998-12-16

<150> 60/113,511

<151> 1998-12-22

<150> 60/115,558

<151> 1999-01-12

<150>60/115,565

<151> 1999-01-12

<150> 60/115,733

<151> 1999-01-12

<150>60/119,341

<151> 1999-02-09

- <151> 2000-03-03
- <150> PCT/US99/12252
- <151> 1999-06-02
- <150> PCT/US99/28634
- <151> 1999-12-01
- <150> PCT/US99/28551
- <151> 1999-12-02
- <150> PCT/US00/03565
- <151> 2000-02-11
- <150> PCT/US00/04414
- <151> 2000-02-22
- <150> PCT/US00/05841
- <151> 2000 -03-02
- <150> PCT/US00/08439
- <151> 2000-03-30
- <150> PCT/US00/14941
- <151> 2000-05-30
- <150> PCT/US00/15264
- <151> 2000-06-02
- <150> PCT/US00/32678
- <151> 2000-12-01
- <140> US 09/866,034
- <141> 2001-05-25
- <160> 38
- <210> 1
- <211> 1283
- <212> DNA
- <213> Homo sapiens
- <400> 1
- cggacgcgtg ggacccatac ttgctggtct gatccatgca caaggcgggg 50
- ctgctaggcc tctgtgcccg ggcttggaat tcggtgcgga tggccagctc 100
- cgggatgacc cgccgggacc cgctcgcaaa taaggtggcc ctggtaacgg 150
- cctccaccga cgggatcggc ttcgccatcg cccggcgttt ggcccaggac 200
- ggggcccatg tggtcgtcag cagccggaag cagcagaatg tggaccaggc 250
- ggtggccacg ctgcaggggg aggggctgag cgtgacgggc accgtgtgcc 300
- atgtggggaa ggcggaggac cgggagcggc tggtggccac ggctgtgaag 350

cttcatqqaq qtatcqatat cctaqtctcc aatgctgctg tcaacccttt 400 ctttqqaaqc ataatqqatq tcactqaqga ggtgtgggac aagactctgg 450 acattaatgt gaaggcccca gccctgatga caaaggcagt ggtgccagaa 500 atggagaaac gaggaggegg eteagtggtg ategtgtett ceatageage 550 cttcaqtcca tctcctggct tcagtcctta caatgtcagt aaaacagcct 600 tgctgggcct gaccaagacc ctggccatag agctggcccc aaggaacatt 650 agggtgaact gcctagcacc tggacttatc aagactagct tcagcaggat 700 getetggatg gacaaggaaa aagaggaaag catgaaagaa accetgegga 750 taagaaggtt aggcgagcca gaggattgtg ctggcatcgt gtctttcctg 800 tgctctgaag atgccagcta catcactggg gaaacagtgg tggtgggtgg 850 aggaaccccg tcccgcctct gaggaccggg agacagccca caggccagag 900 ttgggctcta gctcctggtg ctgttcctgc attcacccac tggcctttcc 950 cacctetget caccttactg ttcacctcat caaatcagtt ctgccctgtg 1000 aaaagateea geetteeetg eegteaaggt ggegtettae tegggattee 1050 tgctgttgtt gtggccttgg gtaaaggcct cccctgagaa cacaggacag 1100 gcctgctgac aaggctgagt ctaccttggc aaagaccaag atattttttc 1150 ctgggccact ggtgaatctg aggggtgatg ggagagaagg aacctggagt 1200 ggaaggagca gagttgcaaa ttaacagctt gcaaatgagg tgcaaataaa 1250 atgcagatga ttgcgcggct ttgaaaaaaa aaa 1283

<210> 2

<211> 278

<212> PRT

<213> Homo sapiens

<400> 2

Met His Lys Ala Gly Leu Leu Gly Leu Cys Ala Arg Ala Trp Asn 1 5 10 15

Ser Val Arg Met Ala Ser Ser Gly Met Thr Arg Arg Asp Pro Leu 20 25 30

Ala Asn Lys Val Ala Leu Val Thr Ala Ser Thr Asp Gly Ile Gly
35 40 45

Phe Ala Ile Ala Arg Arg Leu Ala Gln Asp Gly Ala His Val Val
50 55 60

Val Ser Ser Arg Lys Gln Gln Asn Val Asp Gln Ala Val Ala Thr 65 70 75

Leu Gln Gly	Glu Gly 80	Leu S	er Val	Thr	Gly 85	Thr	Val	Cys	His	Val 90
Gly Lys Ala	Glu Asp 95	Arg G	lu Arg	Leu	Val 100	Ala	Thr	Ala	Val	Lys 105
Leu His Gly	Gly Ile 110	Asp I	le Leu	Val	Ser 115	Asn	Ala	Ala	Val	Asn 120
Pro Phe Phe	Gly Ser 125	Ile M	et Asp	Val	Thr 130	Glu	Glu	Val	Trp	Asp 135
Lys Thr Leu	Asp Ile 140	Asn V	al Lys	Ala	Pro 145	Ala	Leu	Met	Thr	Lys 150
Ala Val Val	Pro Glu 155	Met G	lu Lys	Arg	Gly 160	Gly	Gly	Ser	Val	Val 165
Ile Val Ser	Ser Ile 170	Ala A	la Phe	Ser	Pro 175	Ser	Pro	Gly	Phe	Ser 180
Pro Tyr Asn	Val Ser 185	Lys T	hr Ala	Leu	Leu 190	Gly	Leu	Thr	Lys	Thr 195
Leu Ala Ile	Glu Leu 200	Ala P	ro Arg	Asn	11e 205	Arg	Val	Asn	Cys	Leu 210
Ala Pro Gly	Leu Ile 215	Lys T	hr Ser	Phe	Ser 220	Arg	Met	Leu	Trp	Met 225
Asp Lys Glu	230			-	235					240
Arg Leu Gly	245		-		250					255
Cys Ser Glu	260		-	Thr	Gly 265	Glu	Thr	Val	Val	Val 270
Gly Gly Gly	min m	Com 7	ma Ton							

- <210> 3
- <211> 21
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> Synthetic Oligonucleotide Probe
- <400> 3
- gcataatgga tgtcactgag g 21
- <210> 4
- <211> 23
- <212> DNA
- <213> Artificial Sequence

<220> <223> Synthetic Oligonucleotide Probe <400> 4 agaacaatcc tgctgaaagc tag 23 <210> 5 <211> 46 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 5 gaaacgagga ggcggctcag tggtgatcgt gtcttccata gcagcc 46 <210> 6 <211> 3121 <212> DNA <213> Homo sapiens gegeeetgag eteegeetee gggeeegata geggeatega gagegeetee 50 gtcgaggacc aggcggcgca gggggccggc gggcgaaagg aggatgaggg 100 ggcgcagcag ctgctgaccc tgcagaacca ggtggcgcgg ctggaggagg 150 agaaccgaga ctttctggct gcgctggagg acgccatgga gcagtacaaa 200 ctgcagagcg accggctgcg tgagcagcag gaggagatgg tggaactgcg 250 gctgcggtta gagctggtgc ggccaggctg ggggggcctg cggctcctga 300 atggcetgee teeegggtee titgtgeete gaceteatae ageceeetg 350 gggggtgccc acgcccatgt gctgggcatg gtgccgcctg cctgcctccc 400 tggagatgaa gttggctctg agcagagggg agagcaggtg acaaatggca 450 gggaggetgg agetgagttg etgaetgagg tgaacagget gggaagtgge 500 tetteagetg etteagagga ggaagaggag gaggaggage egeecaggeg 550 gacettacae etgegeagaa ataggateag caactgeagt cagagggegg 600 gggcacgccc agggagtctg ccagagagga agggcccaga gctttgcctt 650 gaggagttgg atgeageeat tecagggtee agageagttg gtgggageaa 700 ggecegagtt caggecegec aggtececcc tgccacagec tcagagtggc 750 ggctggccca ggcccagcag aagatccggg agctggctat caacatccgc 800 atgaaggagg agcttattgg cgagctggtc cgcacaggaa aggcagctca 850

ggccctgaac cgccagcaca gccagcgtat ccgggagctg gagcaggagg 900

cagagcaggt gegggeegag etgagtgaag geeagaggea getgegggag 950 ctcgagggca aggagctcca ggatgctggc gagcggtctc ggctccagga 1000 gttccgcagg agggtcgctg cggcccagag ccaggtgcag gtgctgaagg 1050 agaagaagca ggctacggag cggctggtgt cactgtcggc ccagagtgag 1100 aagcgactgc aggagetega geggaaegtg cageteatge ggeageagea 1150 gggacagetg cagaggegge ttegegagga gaeggageag aageggegee 1200 tggaggcaga aatgagcaag cggcagcacc gcgtcaagga gctggagctg 1250 aagcatgagc aacagcagaa gatcctgaag attaagacgg aagagatcgc 1300 ggccttccag aggaagagge gcagtggcag caacggctct gtggtcagcc 1350 tggaacagca gcagaagatt gaggagcaga agaagtggct ggaccaggag 1400 atggagaagg tgctacagca gcggcgggcg ctggaggagc tggggggagga 1450 gctccacaag cgggaggcca tcctggccaa gaaggaggcc ctgatgcagg 1500 agaagacggg gctggagagc aagcgcctga gatccagcca ggccctcaac 1550 gaggacateg tgcgagtgtc cagccggctg gagcacctgg agaaggagct 1600 gtccgagaag agcgggcagc tgcggcaggg cagcgcccag agccagcagc 1650 agateegegg ggagategae ageetgegee aggagaagga etegetgete 1700 aagcagcgcc tggagatcga cggcaagctg aggcagggga gtctgctgtc 1750 eccegaggag gageggaege tgttecagtt ggatgaggee ategaggeee 1800 tggatgctgc cattgagtat aagaatgagg ccatcacatg ccgccagcgg 1850 gtgetteggg ceteageete gttgetgtee eagtgegaga tgaaceteat 1900 ggccaagete agetacetet eateeteaga gaccagagee eteetetgea 1950 agtattttga caaggtggtg acgctccgag aggagcagca ccagcagcag 2000 attgccttct cggaactgga gatgcagctg gaggagcagc agaggctggt 2050 gtactggetg gaggtggeee tggageggea gegeetggag atggaeegee 2100 agetgaeeet geageagaag gageaegage agaaeatgea getgeteetg 2150 cagcagagte gagaccacet eggtgaaggg ttagcagaca gcaggaggca 2200 gtatgaggcc cggattcaag ctctggagaa ggaactgggc cgttacatgt 2250 ggataaacca ggaactgaaa cagaagctcg gcggtgtgaa cgctgtaggc 2300 cacagcaggg gtggggagaa gaggagectg tgctcggagg gcagacaggc 2350

teetggaaat gaagatgage teeacetgge accegagett etetggetgt 2400 ccccctcac tqaqqqqcc ccccqcaccc qqqaqqaqac qcqqqacttq 2450 gtocacgete egttaceett gacetggaaa egetegagee tgtgtggtga 2500 ggagcagggg tecceegagg aactgaggea gegggaggeg getgageece 2550 tggtggggeg ggtgetteet gtgggtgagg caggeetgee etggaacttt 2600 gggeetttgt ccaageeeg gegggaactg egacgageea geeeggggat 2650 gattgatgtc cggaaaaacc ccctgtaagc cctcggggca gaccctgcct 2700 tggagggaga ctccgagcct gctgaaaggg gcagctgcct gttttgcttc 2750 tgtgaaggge agteettace geacaceeta aateeaggee eteatetgta 2800 ccctcactgg gatcaacaaa tttgggccat ggcccaaaag aactggaccc 2850 tcatttaaca aaataatatg caaattccca ccacttactt ccatgaagct 2900 gtggtaccca attgccqcct tgtgtcttgc tcgaatctca ggacaattct 2950 ggtttcaggc gtaaatggat gtgcttgtag ttcaggggtt tggccaagaa 3000 tcatcacgaa agggtcggtg gcaaccaggt tgtggtttaa atggtcttat 3050 gtatataggg gaaactggga gactttagga tottaaaaaa ccatttaata 3100 aaaaaaaatc tttgaaggga c 3121

<210> 7 <211> 830

<212> PRT

<213> Homo sapiens

<400> 7

Met Glu Gln Tyr Lys Leu Gln Ser Asp Arg Leu Arg Glu Gln Gln 1 5 10 15

Glu Glu Met Val Glu Leu Arg Leu Arg Leu Glu Leu Val Arg Pro
20 25 30

Gly Trp Gly Gly Leu Arg Leu Leu Asn Gly Leu Pro Pro Gly Ser 35 40 45

Phe Val Pro Arg Pro His Thr Ala Pro Leu Gly Gly Ala His Ala 50 55 60

His Val Leu Gly Met Val Pro Pro Ala Cys Leu Pro Gly Asp Glu 65 70 75

Val Gly Ser Glu Gln Arg Gly Glu Gln Val Thr Asn Gly Arg Glu 80 85 90

Ala Gly Ala Glu Leu Leu Thr Glu Val Asn Arg Leu Gly Ser Gly
95 100 105

Ser	Ser	Ala	Ala	Ser 110	Glu	Glu	Glu	Glu	Glu 115	Glu	Glu	Glu	Pro	Pro 120
Arg	Arg	Thr	Leu	His 125	Leu	Arg	Arg	Asn	Arg 130	Ile	Ser	Asn	Суѕ	Ser 135
Gln	Arg	Ala	Gly	Ala 140	Arg	Pro	Gly	Ser	Leu 145	Pro	Glu	Arg	Lys	Gly 150
Pro	Glu	Leu	Cys	Leu 155	Glu	Glu	Leu	Asp	Ala 160	Ala	Ile	Pro	Gly	Ser 165
Arg	Ala	Val	Gly	Gly 170	Ser	Lys	Ala	Arg	Val 175	Gln	Ala	Arg	Gln	Val 180
Pro	Pro	Ala	Thr	Ala 185	Ser	Glu	Trp	Arg	Leu 190	Ala	Gln	Ala	Gln	Gln 195
Lys	Ile	Arg	Glu	Leu 200	Ala	Ile	Asn	Ile	Arg 205	Met	Lys	Glu	Glu	Leu 210
Ile	Gly	Glu	Leu	Val 215	Arg	Thr	Gly	Lys	Ala 220	Ala	Gln	Ala	Leu	Asn 225
Arg	Gln	His	Ser	Gln 230	Arg	Ile	Arg	Glu	Leu 235	Glu	Gln	Glu	Ala	Glu 240
Gln	Val	Arg	Ala	Glu 245	Leu	Ser	Glu	Gly	Gln 250	Arg	Gln	Leu	Arg	Glu 255
Leu	Glu	Gly	Lys	Glu 260	Leu	Gln	Asp	Ala	Gly 265	Glu	Arg	Ser	Arg	Leu 270
Gln	Glu	Phe	Arg	Arg 275	Arg	Val	Ala	Ala	Ala 280	Gln	Ser	Gln	Val	Gln 285
Val	Leu	Lys	Glu	Lys 290	Lys	Gln	Ala	Thr	Glu 295	Arg	Leu	Val	Ser	Leu 300
Ser	Ala	Gln	Ser	Glu 305	Lys	Arg	Leu	Gln	Glu 310	Leu	Glu	Arg	Asn	Val 315
Gln	Leu	Met	Arg	Gln 320	Gln	Gln	Gly	Gln	Leu 325	Gln	Arg	Arg	Leu	Arg 330
Glu	Glu	Thr	Glu	Gln 335	Lys	Arg	Arg	Leu	Glu 340	Ala	Glu	Met	Ser	Lys 345
Arg	Gln	His	Arg	Val 350	Lys	Glu	Leu	Glu	Leu 355	Lys	His	Glu	Gln	Gln 360
Gln	Lys	Ile	Leu	Lys 365	Ile	Lys	Thr	Glu	Glu 370	Ile	Ala	Ala	Phe	Gln 375
Arg	Lys	Arg	Arg	Ser 380	Gly	Ser	Asn	Gly	Ser 385	Val	Val	Ser	Leu	Glu 390
Gln	Gln	Gln	Lys	Ile	Glu	Glu	Gln	Lys	Lys	Trp	Leu	Asp	Gln	Glu

				395					400					405
Met	Glu	Lys	Val	Leu 410	Gln	Gln	Arg	Arg	Ala 415	Leu	Glu	Glu	Leu	Gly 420
Glu	Glu	Leu	His	Lys 425	Arg	Glu	Ala	Ile	Leu 430	Ala	Lys	Lys	Glu	Ala 435
Leu	Met	Gln	Glu	Lys 440	Thr	Gly	Leu	Glu	Ser 445	Lys	Arg	Leu	Arg	Ser 450
Ser	Gln	Ala	Leu	Asn 455	Glu	Asp	Ile	Val	Arg 460	Val	Ser	Ser	Arg	Leu 465
Glu	His	Leu	Glu	Lys 470	Glu	Leu	Ser	Glu	Lys 475	Ser	Gly	Gln	Leu	Arg 480
Gln	Gly	Ser	Ala	Gln 485	Ser	Gln	Gln	Gln	Ile 490	Arg	Gly	Glu	Ile	Asp 495
Ser	Leu	Arg	Gln	Glu 500	Lys	Asp	Ser	Leu	Leu 505	Lys	Gln	Arg	Leu	Glu 510
Ile	Asp	Gly	Lys	Leu 515	Arg	Gln	Gly	Ser	Leu 520	Leu	Ser	Pro	Glu	Glu 525
Glu	Arg	Thr	Leu	Phe 530	Gln	Leu	Asp	Glu	Ala 535	Ile	Glu	Ala	Leu	Asp 540
Ala	Ala	Ile	Glu	Tyr 545	Lys	Asn	Glu	Ala	Ile 550	Thr	Cys	Arg	Gln	Arg 555
Val	Leu	Arg	Ala	Ser 560	Ala	Ser	Leu	Leu	Ser 565	Gln	Cys	Glu	Met	Asn 570
Leu	Met	Ala	Lys	Leu 575	Ser	Tyr	Leu	Ser	Ser 580	Ser	Glu	Thr	Arg	Ala 585
Leu	Leu	Cys	Lys	Tyr 590	Phe	Asp	Lys	Val	Val 595	Thr	Leu	Arg	Glu	Glu 600
Gln	His	Gln	Gln	Gln 605	Ile	Ala	Phe	Ser	Glu 610	Leu	Glu	Met	Gln	Leu 615
Glu	Glu	Gln	Gln	Arg 620	Leu	Val	Tyr	Trp	Leu 625	Glu	Val	Ala	Leu	Glu 630
Arg	Gln	Arg	Leu	Glu 635	Met	Asp	Arg	Gln	Leu 640	Thr	Leu	Gln	Gln	Lys 645
Glu	His	Glu	Gln	Asn 650	Met	Gln	Leu	Leu	Leu 655	Gln	Gln	Ser	Arg	Asp 660
His	Leu	Gly	Glu	Gly 665	Leu	Ala	Asp	Ser	Arg 670	Arg	Gln	Tyr	Glu	Ala 675
Arg	Ile	Gln	Ala	Leu 680	Glu	Lys	Glu	Leu	Gly 685	Arg	Tyr	Met	Trp	Ile 690

 Asn
 Gln
 Glu
 Leu
 Lys
 Gln
 Lys
 Leu
 Gly
 Val
 Asn
 Ala
 Val
 705

 His
 Ser
 Arg
 Gly
 Gly
 Glu
 Lys
 Arg
 Ser
 Leu
 Cys
 Ser
 Glu
 Gly
 Arg
 720

 Gln
 Ala
 Pro
 Gly
 Asn
 Glu
 Asp
 Glu
 Leu
 His
 Leu
 His
 Leu
 Ala
 Pro
 Glu
 735

 Leu
 Trp
 Leu
 Ser
 Pro
 Leu
 Thr
 Glu
 Gly
 Ala
 Pro
 Arg
 Thr
 Arg
 Glu
 735

 Glu
 Thr
 Arg
 Asp
 Leu
 Thr
 Glu
 Gly
 Ala
 Pro
 Leu
 Pro
 Arg
 Pro
 Leu
 Pro
 Glu
 Pro
 Pro

<210> 8 <211> 662 <212> DNA <213> Homo sapiens

attetectag ageatettig gaageatgag geeaegatge tigeatettigg 50 etettigtetig etggataaca gitetteetee teeagitgite aaaaggaaet 100 acagaegete etgitiggete aggaetigigg etgitigeeage egacaeceag 150 gitigggaae aagatetaca accetteaga geagitgetig tatgatgatig 200 ecatettate ettaaaggag accegeeget gitiggeteeae etgeaecette 250 tiggeeetiget tigagetetig etgiteeegag tetititiggee eeaageagaa 300 gitiettigig aagitigaggi tiettigggaae teetigaaeetia 350 eteecatete eeggagetigi accaggaaea ggaggeaegi eetgitaeeea 400 taaaaaecee aggeteeaet gigeagaegi agaaeaggig agaagaaegi 500 gaetteagag gaatggetea atgaeatage titiggagag gaageaeeti 500 gaetteagag ggatggetea atgaeatage titiggagag ageeeagetig 550

gggatggcca gaetteaggg gaagaatgee tteetgette ateceettte 600 cageteecet teeegetgag agecaettte ateggeaata aaateeecea 650 catttaceat et 662

<210> 9

<211> 125

<212> PRT

<213> Homo sapiens

<400> 9

Met Arg Pro Arg Cys Cys Ile Leu Ala Leu Val Cys Trp Ile Thr 1 5 10 15

Val Phe Leu Leu Gln Cys Ser Lys Gly Thr Thr Asp Ala Pro Val 20 25 30

Gly Ser Gly Leu Trp Leu Cys Gln Pro Thr Pro Arg Cys Gly Asn 35 40 45

Lys Ile Tyr Asn Pro Ser Glu Gln Cys Cys Tyr Asp Asp Ala Ile 50 55 60

Leu Ser Leu Lys Glu Thr Arg Arg Cys Gly Ser Thr Cys Thr Phe
65 70 75

Trp Pro Cys Phe Glu Leu Cys Cys Pro Glu Ser Phe Gly Pro Gln 80 85 90

Gln Lys Phe Leu Val Lys Leu Arg Val Leu Gly Met Lys Ser Gln 95 100 105

Cys His Leu Ser Pro Ile Ser Arg Ser Cys Thr Arg Asn Arg Arg 110 115 120

His Val Leu Tyr Pro

<210> 10

<211> 1942

<212> DNA

<213> Homo sapiens

<400> 10
 cccaegegte cgccaegeg tecgggtgec actegegege cggccgeget 50
 ccgggettet cttttecete cgaegegea cggetgeca gaeatteegg 100
 ctgccgggte tggagagete cecgaaceee tecgeggaga ggagegagge 150
 ggegecaggg tggccecegg ggegegettg gteteggaga agegggaeg 200
 aggeeggagg atgagegaet gagggegaeg egggaettga egegagttgg 250
 ggeegegaet aceggeaget gaeagegga tgagegaete eccagagaeg 300
 ccctageeeg gtgtgegege caggeggage gegeaggtgg ggetgggetg 350

ttagtggtee geeceaegeg ggtegeegge eggeecagga tgggegetgg 400 caaccoggge cogogocogo ogotgetado cotgogocog etgogagedo 450 ggegteegge eegegeetg egeteatgga eggeggetee eggetggegg 500 cggcgcgccc ccgggctgtg aatgcgactc gcccctcggc cgcgctcccc 550 gecegecege eegeegggae gtggtagggg atgeceaget ecaetgegat 600 ggcagttggc gcgctctcca gttccctcct ggtcacctgc tgcctgatgg 650 tggctctgtg cagtccgagc atcccgctgg agaagctggc ccaggcacca 700 gagcagcegg gecaggagaa gegtgageae gecaeteggg aeggeeeggg 750 gegggtgaac gageteggge geeeggegag ggaegaggge ggeageggee 800 gggactggaa gagcaagagc ggccgtgggc tcgccggccg tgagccgtgg 850 agcaagctga agcaggcctg ggtctcccag ggcgggggg ccaaggccgg 900 ggatctgcag gtccggcccc gcggggacac cccgcaggcg gaagccctgg 950 ccgcagccgc ccaggacgcg attggcccgg aactcgcgcc cacgcccgag 1000 ccacccgagg agtacgtgta cccggactac cgtggcaagg gctgcgtgga 1050 cgagagcggc ttcgtgtacg cgatcgggga gaagttcgcg ccgggcccct 1100 eggeetgeee gtgeetgtge acegaggagg ggeegetgtg egegeageee 1150 gagtgcccga ggctgcaccc gcgctgcatc cacgtcgaca cgagccagtg 1200 ctgcccgcag tgcaaggaga ggaagaacta ctgcgagttc cggggcaaga 1250 cctatcagac tttggaggag ttcgtggtgt ctccatgcga gaggtgtcgc 1300 tgtgaagcca acggtgaggt gctatgcaca gtgtcagcgt gtccccagac 1350 ggagtgtgtg gaccctgtgt acgagcctga tcagtgctgt cccatctgca 1400 aaaatggtcc aaactgcttt gcagaaaccg cggtgatccc tgctggcaga 1450 gaagtgaaga ctgacgagtg caccatatgc cactgtactt atgaggaagg 1500 cacatggaga atcgagcggc aggccatgtg cacgagacat gaatgcaggc 1550 aaatgtagac getteecaga acacaaacte tgaettttte tagaacattt 1600 tactgatgtg aacattctag atgactctgg gaactatcag tcaaagaaga 1650 cttttgatga ggaataatgg aaaattgttg gtacttttcc ttttcttgat 1700 aacagttact acaacagaag gaaatggata tatttcaaaa catcaacaag 1750 aactttgggc ataaaatcct tctctaaata aatgtgctat tttcacagta 1800

<210> 11

<211> 325

<212> PRT

<213> Homo sapiens

<400> 11

Met Pro Ser Ser Thr Ala Met Ala Val Gly Ala Leu Ser Ser Ser 1 5 10 15

Leu Leu Val Thr Cys Cys Leu Met Val Ala Leu Cys Ser Pro Ser 20 25 30

Ile Pro Leu Glu Lys Leu Ala Gln Ala Pro Glu Gln Pro Gly Gln
35 40 45

Glu Lys Arg Glu His Ala Thr Arg Asp Gly Pro Gly Arg Val Asn
50 55 60

Glu Leu Gly Arg Pro Ala Arg Asp Glu Gly Gly Ser Gly Arg Asp
65 70 75

Trp Lys Ser Lys Ser Gly Arg Gly Leu Ala Gly Arg Glu Pro Trp 80 85 90

Ser Lys Leu Lys Gln Ala Trp Val Ser Gln Gly Gly Ala Lys 95 100 105

Ala Gly Asp Leu Gln Val Arg Pro Arg Gly Asp Thr Pro Gln Ala 110 115 120

Glu Ala Leu Ala Ala Ala Gln Asp Ala Ile Gly Pro Glu Leu 125 130 135

Ala Pro Thr Pro Glu Pro Pro Glu Glu Tyr Val Tyr Pro Asp Tyr
140 145 150

Arg Gly Lys Gly Cys Val Asp Glu Ser Gly Phe Val Tyr Ala Ile 155 160 165

Gly Glu Lys Phe Ala Pro Gly Pro Ser Ala Cys Pro Cys Leu Cys 170 175 180

Thr Glu Glu Gly Pro Leu Cys Ala Gln Pro Glu Cys Pro Arg Leu 185 190 195

His Pro Arg Cys Ile His Val Asp Thr Ser Gln Cys Cys Pro Gln 200 205 210

Cys Lys Glu Arg Lys Asn Tyr Cys Glu Phe Arg Gly Lys Thr Tyr 215 220 225

Gln Thr Leu Glu Glu Phe Val Val Ser Pro Cys Glu Arg Cys Arg

i mile
i com
1000
127
The state of
4;
ļazis
man man Huli judi maj maj
man man Huli judi maj maj

230 235 240 Cys Glu Ala Asn Gly Glu Val Leu Cys Thr Val Ser Ala Cys Pro Gln Thr Glu Cys Val Asp Pro Val Tyr Glu Pro Asp Gln Cys Cys 260 Pro Ile Cys Lys Asn Gly Pro Asn Cys Phe Ala Glu Thr Ala Val 275 280 Ile Pro Ala Gly Arg Glu Val Lys Thr Asp Glu Cys Thr Ile Cys His Cys Thr Tyr Glu Glu Gly Thr Trp Arg Ile Glu Arg Gln Ala Met Cys Thr Arg His Glu Cys Arg Gln Met 320 <210> 12 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 12 gaggtgtcgc tgtgaagcca acgg 24 <210> 13 <211> 24 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 13 cgctcgattc tccatgtgcc ttcc 24 <210> 14 <211> 45 <212> DNA <213> Artificial Sequence <220> <223> Synthetic Oligonucleotide Probe <400> 14 gacggagtgt gtggaccctg tgtacgagcc tgatcagtgc tgtcc 45 <210> 15 <211> 1587 <212> DNA <213> Homo sapiens

<400> 15 cagecacaga egggteatga gegeggtatt aetgetggee eteetggggt 50 teatectece actgecagga gtgeaggege tgetetgeca gtttgggaca 100 gttcagcatg tgtggaaggt gtccgaccta ccccggcaat ggacccctaa 150 gaacaccage tgcgacageg gettggggtg ccaggacaeg ttgatgetca 200 ttgagagegg acceeaagtg ageetggtge tetecaaggg etgeaeggag 250 gccaaggacc aggageceeg egteaetgag caeeggatgg gcceeggeet 300 ctccctgatc tcctacacct tcgtgtgccg ccaggaggac ttctgcaaca 350 acctegttaa eteceteeeg etttgggeee caeageeece ageagaeeca 400 ggatcettga ggtgeccagt etgettgtet atggaagget gtetggaggg 450 gacaacagaa gagatetgee ecaaggggae cacacactgt tatgatggee 500 tootoaggot caggggagga ggcatottot ccaatotgag agtocaggga 550 tgcatgcccc agccaggttg caacctgctc aatgggacac aggaaattgg 600 gcccgtgggt atgactgaga actgcaatag gaaagatttt ctgacctgtc 650 ategggggae caccattatg acacaeggaa aettggetea agaaeeeaet 700 gattggacca catcgaatac cgagatgtgc gaggtggggc aggtgtgtca 750 ggagacgetg etgeteatag atgtaggaet cacateaace etggtgggga 800 caaaaggetg cagcactgtt ggggeteaaa atteecagaa gaccaccate 850 cactcagece etectggggt gettgtggee tectatacee acttetgete 900 ctcggacctg tgcaatagtg ccagcagcag cagcgttctg ctgaactccc 950 tecetectea agetgeeest gteecaggag aceggeagtg tectacetgt 1000 gtgcagcccc ttggaacctg ttcaagtggc tccccccgaa tgacctgccc 1050 caggggggcc actcattgtt atgatgggta cattcatctc tcaggaggtg 1100 ggctgtccac caaaatgagc attcagggct gcgtggccca accttccagc 1150 ttottgttga accacaccag acaaatcggg atottetetg cgcgtgagaa 1200 qcqtqatqtg cagcctcctg cctctcagca tgagggaggt ggggctgagg 1250 geetggagte teteaettgg ggggtgggge tggeaetgge cecagegetg 1300 tggtggggag tggtttgccc ttcctgctaa ctctattacc cccacgattc 1350 ttcaccgctg ctgaccaccc acactcaacc tccctctgac ctcataacct 1400 aatggccttg gacaccagat tettteeeat tetgteeatg aateatette 1450 cccacacac atcattcata tetactcace taacagcaac actggggaga 1500 geetggagca teeggacttg ceetatggga gaggggacge tggaggagtg 1550 getgcatgta tetgataata cagaceetgt cetttea 1587

<210> 16

<211> 437

<212> PRT

<213> Homo sapiens

<400> 16

Met Ser Ala Val Leu Leu Leu Ala Leu Leu Gly Phe Ile Leu Pro 1 5 10 15

Leu Pro Gly Val Gln Ala Leu Cys Gln Phe Gly Thr Val Gln
20 25 30

His Val Trp Lys Val Ser Asp Leu Pro Arg Gln Trp Thr Pro Lys
35 40 45

Asn Thr Ser Cys Asp Ser Gly Leu Gly Cys Gln Asp Thr Leu Met
50 55 60

Leu Ile Glu Ser Gly Pro Gln Val Ser Leu Val Leu Ser Lys Gly
65 70 75

Cys Thr Glu Ala Lys Asp Gln Glu Pro Arg Val Thr Glu His Arg 80 85 90

Met Gly Pro Gly Leu Ser Leu Ile Ser Tyr Thr Phe Val Cys Arg 95 100 105

Gln Glu Asp Phe Cys Asn Asn Leu Val Asn Ser Leu Pro Leu Trp
110 115 120

Ala Pro Gln Pro Pro Ala Asp Pro Gly Ser Leu Arg Cys Pro Val 125 130 135

Cys Leu Ser Met Glu Gly Cys Leu Glu Gly Thr Thr Glu Glu Ile 140 145 150

Cys Pro Lys Gly Thr Thr His Cys Tyr Asp Gly Leu Leu Arg Leu 155 160 165

Arg Gly Gly Gle Phe Ser Asn Leu Arg Val Gln Gly Cys Met
170 175 180

Pro Gln Pro Gly Cys Asn Leu Leu Asn Gly Thr Gln Glu Ile Gly 185 190 195

Pro Val Gly Met Thr Glu Asn Cys Asn Arg Lys Asp Phe Leu Thr 200 205 210

Cys His Arg Gly Thr Thr Ile Met Thr His Gly Asn Leu Ala Gln
215 220 225

Glu Pro Thr Asp Trp Thr Thr Ser Asn Thr Glu Met Cys Glu Val

				230					235					240
Gly	Gln	Val	Cys	Gln 245	Glu	Thr	Leu	Leu	Leu 250	Ile	Asp	Val	Gly	Leu 255
Thr	Ser	Thr	Leu	Val 260	Gly	Thr	Lys	Gly	Cys 265	Ser	Thr	Val	Gly	Ala 270
Gln	Asn	Ser	Gln	Lys 275	Thr	Thr	Ile	His	Ser 280	Ala	Pro	Pro	Gly	Val 285
Leu	Val	Ala	Ser	Tyr 290	Thr	His	Phe	Cys	Ser 295	Ser	Asp	Leu	Cys	Asn 300
Ser	Ala	Ser	Ser	Ser 305	Ser	Val	Leu	Leu	Asn 310	Ser	Leu	Pro	Pro	Gln 315
Ala	Ala	Pro	Val	Pro 320	Gly	Asp	Arg	Gln	Cys 325	Pro	Thr	Cys	Val	Gln 330
Pro	Leu	Gly	Thr	Cys 335	Ser	Ser	Gly	Ser	Pro 340	Arg	Met	Thr	Cys	Pro 345
Arg	Gly	Ala	Thr	His 350	Сув	Tyr	Asp	Gly	Tyr 355	Ile	His	Leu	Ser	Gly 360
Gly	Gly	Leu	Ser	Thr 365	Lys	Met	Ser	Ile	Gln 370	Gly	Сув	Val	Ala	Gln 375
Pro	Ser	Ser	Phe	Leu 380	Leu	Asn	His	Thr	Arg 385	Gln	Ile	Gly	Ile	Phe 390
Ser	Ala	Arg	Glu	Lys 395	Arg	Asp	Val	Gln	Pro 400	Pro	Ala	Ser	Gln	His 405
Glu	Gly	Gly	Gly	Ala 410	Glu	Gly	Leu	Glu	Ser 415	Leu	Thr	Trp	Gly	Val 420
-		Ala	Leu	Ala 425	Pro	Ala	Leu	Trp	Trp 430	Gly	Val	Val	Cya	Pro 435
Car	Cvc													

Ser Cys

<210> 17

<211> 2387

<212> DNA

<213> Homo sapiens

<400> 17

egacgatget aegegegee ggetgeetee teeggaeete egtagegee 50
geegeggeee tggetgegge getgeteteg tegettgege getgetetet 100
tetagageeg agggaeeegg tggeetegte geteageeee tattteggea 150
eeaagaeteg etaegaggat gteaaeeeeg tgetattgte gggeeeegag 200

gctccgtggc gggaccctga gctgctggag gggacctgca ccccggtgca 250 gctggtcgcc ctcattcgcc acggcacccg ctaccccacg gtcaaacaga 300 teegeaaget gaggeagetg caegggttge tgeaggeeeg egggteeagg 350 gatggcgggg ctagtagtac cggcagccgc gacctgggtg cagcgctggc 400 cgactggcct ttgtggtacg cggactggat ggacgggcag ctagtagaga 450 agggacggca ggatatgcga cagctggcgc tgcgtctggc ctcgctcttc 500 ceggeeettt teageegtga gaactaegge egeetgegge teateaceag 550 ttccaagcac cgctgcatgg atagcagcgc cgccttcctg caggggctgt 600 ggcagcacta ccaccetgge ttgccgccgc cggacgtcgc agatatggag 650 tttggacctc caacagttaa tgataaacta atgagatttt ttgatcactg 700 tgagaagttt ttaactgaag tagaaaaaaa tgctacagct ctttatcacg 750 tggaagcett caaaactgga ccagaaatgc agaacatttt aaaaaaagtt 800 gcagctactt tgcaagtgcc agtaaatgat ttaaatgcag atttaattca 850 agtageettt tteacetgtt catttgaeet ggeaattaaa ggtgttaaat 900 ctccttggtg tgatgttttt gacatagatg atgcaaaggt attagaatat 950 ttaaatgate tgaaacaata ttggaaaaga ggatatgggt atactattaa 1000 cagtegatee agetgeacet tgttteagga tatettteag caettggaca 1050 aagcagttga acagaaacaa aggteteage caatttette tecagteate 1100 ctccagtttg gtcatgcaga gactettett ccactgettt ctctcatggg 1150 ctacttcaaa gacaaggaac ccctaacagc gtacaattac aaaaaacaaa 1200 tgcatcggaa gttccgaagt ggtctcattg taccttatgc ctcgaacctg 1250 atatttgtgc tttaccactg tgaaaatgct aagactccta aagaacaatt 1300 ccgagtgcag atgttattaa atgaaaaggt gttacctttg gcttactcac 1350 aagaaactgt ttcattttat gaagatctga agaaccacta caaggacatc 1400 cttcagagtt gtcaaaccag tgaagaatgt gaattagcaa gggctaacag 1450 tacatotgat gaactatgag taactgaaga acatttttaa ttotttagga 1500 atotgoaatg agtgattaca tgottgtaat aggtaggcaa ttoottgatt 1550 acaggaaget tttatattae ttgagtattt etgtetttte acagaaaaac 1600 attgggtttc tctctgggtt tggacatgaa atgtaagaaa agatttttca 1650 ceetgaagcage tetettaagg agaaacaaat etatttagag aaacagetgg 1700 ceetgcaaat gtttacagaa atgaaattet teetacttat ataagaaate 1750 teaccactgag atagaattgt gattteataa taacacttga aaagtgetgg 1800 agtaacaaaa tateteagtt ggaccateet taacettgat gaactgteta 1850 ggaactttae agattgttet geagttetet ettetttee teaggtagga 1900 cagetetage attteetaa teaggaatat tgtgggtaage tgggagtate 1950 actetggaag aaagtaacat etecagatga gaatttgaaa eaagaacaag 2000 agtgttgtaa aaggacacet teacetgaage aagteggaaa gtacaatgaa 2050 aataaatatt tettggtattt atttatgaaa tatttgaaca tttttteaat 2100 aatteetttt tacttetagg aagteteaaa agaccatett aaattattat 2150 atgtttggae aattageaae tttteattet gteacttgge teegatttte 2250 ataatteet attattgaa atgtatett tggttgtttg atttteett 2250 cetataataaa gaaaattett gagttettaa aaatageag aaagaacace taaatageag aagtetetta aaattattat 2350 ettettettg aaatagetet gagttettga aaatgeegtg aaagtatttg 2350 ettaaataaa gaaaattett gtggaetttaa aaaaaaa 2387

<210> 18

<211> 487

<212> PRT

<213> Homo sapiens

<400> 18

Met Leu Arg Ala Pro Gly Cys Leu Leu Arg Thr Ser Val Ala Pro 1 5 10 15

Ala Ala Leu Ala Ala Leu Leu Ser Ser Leu Ala Arg Cys 20 25 30

Ser Leu Leu Glu Pro Arg Asp Pro Val Ala Ser Ser Leu Ser Pro 35 40 45

Tyr Phe Gly Thr Lys Thr Arg Tyr Glu Asp Val Asn Pro Val Leu
50 55 60

Leu Ser Gly Pro Glu Ala Pro Trp Arg Asp Pro Glu Leu Leu Glu 65 70 75

Gly Thr Cys Thr Pro Val Gln Leu Val Ala Leu Ile Arg His Gly 80 85 90

Thr Arg Tyr Pro Thr Val Lys Gln Ile Arg Lys Leu Arg Gln Leu 95 100 105

His Gly Leu Leu Gln Ala Arg Gly Ser Arg Asp Gly Gly Ala Ser

				110					115					120
Ser	Thr	Gly	Ser	Arg 125	Asp	Leu	Gly	Ala	Ala 130	Leu	Ala	Asp	Trp	Pro 135
Leu	Trp	Tyr	Ala	Asp 140	Trp	Met	Asp	Gly	Gln 145	Leu	Val	Glu	Lys	Gly 150
Arg	Gln	Asp	Met	Arg 155	Gln	Leu	Ala	Leu	Arg 160	Leu	Ala	Ser	Leu	Phe 165
Pro	Ala	Leu	Phe	Ser 170	Arg	Glu	Asn	Tyr	Gly 175	Arg	Leu	Arg	Leu	Ile 180
Thr	Ser	Ser	Lys	His 185	Arg	Cys	Met	Asp	Ser 190	Ser	Ala	Ala	Phe	Leu 195
Gln	Gly	Leu	Trp	Gln 200	His	Tyr	His	Pro	Gly 205	Leu	Pro	Pro	Pro	Asp 210
Val	Ala	Asp	Met	Glu 215	Phe	Gly	Pro	Pro	Thr 220	Val	Asn	Asp	Lys	Leu 225
Met	Arg	Phe	Phe	Asp 230	His	Cys	Glu	Lys	Phe 235	Leu	Thr	Glu	Val	Glu 240
Lys	Asn	Ala	Thr	Ala 245	Leu	Tyr	His	Val	Glu 250	Ala	Phe	Lys	Thr	Gly 255
Pro	Glu	Met	Gln	Asn 260	Ile	Leu	Lys	Lys	Val 265	Ala	Ala	Thr	Leu	Gln 270
Val	Pro	Val	Asn	Asp 275	Leu	Asn	Ala	Asp	Leu 280	Ile	Gln	Val	Ala	Phe 285
Phe	Thr	Cys	Ser	Phe 290	Asp	Leu	Ala	Ile	Lys 295	Gly	Val	Lys	Ser	Pro 300
Trp	Cys	Asp	Val	Phe 305	Asp	Ile	Asp	Asp	Ala 310	Lys	Val	Leu	Glu	Tyr 315
Leu	Asn	Asp	Leu	Lys 320	Gln	Tyr	Trp	Lys	Arg 325	Gly	Tyr	Gly	Tyr	Thr 330
Ile	Asn	Ser	Arg	Ser 335	Ser	Cys	Thr	Leu	Phe 340	Gln	Asp	Ile	Phe	Gln 345
His	Leu	Asp	Lys	Ala 350	Val	Glu	Gln	Lys	Gln 355	Arg	Ser	Gln	Pro	Ile 360
Ser	Ser	Pro	Val	Ile 365	Leu	Gln	Phe	Gly	His 370	Ala	Glu	Thr	Leu	Leu 375
Pro	Leu	Leu	Ser	Leu 380	Met	Gly	Tyr	Phe	Lys 385	Asp	Lys	Glu	Pro	Leu 390
Thr	Ala	Tyr	Asn	Tyr 395	Lys	Lys	Gln	Met	His 400	Arg	Lys	Phe	Arg	Ser 405

Gly Leu Ile Val Pro Tyr Ala Ser Asn Leu Ile Phe Val Leu Tyr 420

His Cys Glu Asn Ala Lys Thr Pro Lys Glu Gln Phe Arg Val Gln 435

Met Leu Leu Asn Glu Lys Val Leu Pro Leu Ala Tyr Ser Gln Glu 440

Thr Val Ser Phe Tyr Glu Asp Leu Lys Asn His Tyr Lys Asp Ile 465

Leu Gln Ser Cys Gln Thr Ser Glu Glu Cys Glu Leu Ala Arg Ala 480

Asn Ser Thr Ser Asp Glu Leu

Asn Ser Thr Ser Asp Glu Leu 485

<210> 19

<211> 3554

<212> DNA

<213> Homo sapiens

<400> 19 gggactacaa geegegeege getgeegetg geeceteage aaccetegae 50 atqqcqctqa qqcqqccacc qcqactccgg ctctgcgctc ggctgcctga 100 cttcttcctg ctgctgcttt tcaggggctg cctgataggg gctgtaaatc 150 tcaaatccag caatcgaacc ccagtggtac aggaatttga aagtgtggaa 200 ctgtcttgca tcattacgga ttcgcagaca agtgacccca ggatcgagtg 250 gaagaaaatt caagatgaac aaaccacata tgtgtttttt gacaacaaaa 300 ttcagggaga cttggcgggt cgtgcagaaa tactggggaa gacatccctg 350 aagatetgga atgtgacaeg gagagaetea gecetttate getgtgaggt 400 cqttqctcga aatqaccgca aggaaattga tgagattgtg atcgagttaa 450 ctgtgcaagt gaagccagtg accectgtct gtagagtgcc gaaggctgta 500 ccagtaggca agatggcaac actgcactgc caggagagtg agggccaccc 550 ceggeeteae tacagetggt ategeaatga tgtaccaetg cecaeggatt 600 ccagagecaa teccagattt egeaattett ettteeaett aaactetgaa 650 acaggcactt tggtgttcac tgctgttcac aaggacgact ctgggcagta 700 ctactgcatt gcttccaatg acgcaggete agccaggtgt gaggagcagg 750 agatggaagt ctatgacctg aacattggcg gaattattgg gggggttctg 800 gttgtccttg ctgtactggc cctgatcacg ttgggcatct gctgtgcata 850

cagacgtggc tacttcatca acaataaaca ggatggagaa agttacaaga 900 acccagggaa accagatgga gttaactaca teegeactga egaggaggge 950 gacttcagae acaagtcate gtttgtgate tgagaeeege ggtgtggetg 1000 agagegeaca gagegeacgt geacatacet etgetagaaa eteetgteaa 1050 ggcagcgaga gctgatgcac tcggacagag ctagacactc attcagaagc 1100 ttttegtttt ggecaaagtt gaccactact ettettaete taacaageca 1150 catgaataga agaattttcc tcaagatgga cccggtaaat ataaccacaa 1200 ggaagegaaa etgggtgegt teaetgagtt gggtteetaa tetgtttetg 1250 geotgattee egeatgagta ttagggtgat ettaaagagt ttgeteaegt 1300 aaacgcccgt gctgggccct gtgaagccag catgttcacc actggtcgtt 1350 cagcagccac gacagcacca tgtgagatgg cgaggtggct ggacagcacc 1400 agcagegeat eeeggeggga aeecagaaaa ggettettae aeagcageet 1450 tacttcatcg gcccacagac accacegeag tttettetta aaggetetge 1500 tgateggtgt tgeagtgtee attgtggaga agetttttgg ateageattt 1550 tgtaaaaaca accaaaatca ggaaggtaaa ttggttgctg gaagagggat 1600 cttgcctgag gaaccctgct tgtccaacag ggtgtcagga tttaaggaaa 1650 accttegtet taggetaagt etgaaatggt actgaaatat gettttetat 1700 gggtcttgtt tattttataa aattttacat ctaaattttt gctaaggatg 1750 tattttgatt attgaaaaga aaatttetat ttaaaetgta aatatattgt 1800 catacaatgt taaataacct atttttttaa aaaagttcaa cttaaggtag 1850 aagttccaag ctactagtgt taaattggaa aatatcaata attaagagta 1900 ttttacccaa ggaatcctct catggaagtt tactgtgatg ttccttttct 1950 cacacaagtt ttagcctttt tcacaaggga actcatactg tctacacatc 2000 agaccatagt tgcttaggaa acctttaaaa attccagtta agcaatgttg 2050 aaateagttt geatetette aaaagaaace teteaggtta getttgaact 2100 geotetteet gagatgacta ggacagtetg tacccagagg ccacccagaa 2150 gccctcagat gtacatacac agatgccagt cagctcctgg ggttgcgcca 2200 ggegeeeeg etetagetea etgttgeete getgtetgee aggaggeeet 2250 gccatcottg ggccctggca gtggctgtgt cccagtgagc tttactcacg 2300

tggccettge tteatecage acagetetea ggtgggcaet geagggaeae 2350 tggtgtcttc catgtagcgt cccagctttg ggctcctgta acagacctct 2400 ttttggttat ggatggctca caaaataggg cccccaatgc tattttttt 2450 ttttaagttt gtttaattat ttgttaagat tgtctaaggc caaaggcaat 2500 tgcgaaatca agtctgtcaa gtacaataac attttaaaaa gaaaatggat 2550 cocactgtte etetttgeca cagagaaage acccagaege cacaggetet 2600 gtcgcatttc aaaacaaacc atgatggagt ggcggccagt ccagcctttt 2650 aaagaacgtc aggtggagca gccaggtgaa aggcctggcg gggaggaaag 2700 tgaaacgcct gaatcaaaag cagttttcta attttgactt taaatttttc 2750 atcogoogga gacactgoto coatttgtgg ggggacatta gcaacatcac 2800 teagaageet gtgttettea agageaggtg tteteageet cacatgeeet 2850 gccgtgctgg actcaggact gaagtgctgt aaagcaagga gctgctgaga 2900 aggagcaete cactgtgtge etggagaatg geteteacta eteacettgt 2950 ettteagett eeagtgtett gggtttttta taetttgaea gettttttt 3000 aattgeatae atgagaetgt gttgaetttt tttagttatg tgaaacaett 3050 tgccgcaggc cgcctggcag aggcaggaaa tgctccagca gtggctcagt 3100 getecetggt gtetgetgea tggeateetg gatgettage atgeaagtte 3150 cctccatcat tgccaccttg gtagagaggg atggctcccc accctcageg 3200 ttggggattc acgetecage etcettettg gttgtcatag tgatagggta 3250 gcottattgc cocctottot tataccotaa aacottotac actagtgcca 3300 tgggaaccag gtctgaaaaa gtagagagaa gtgaaagtag agtctgggaa 3350 gtagctgcct ataactgaga ctagacggaa aaggaatact cgtgtatttt 3400 aagatatgaa tgtgactcaa gactcgaggc cgatacgagg ctgtgattct 3450 gcctttggat ggatgttgct gtacacagat gctacagact tgtactaaca 3500 caccgtaatt tggcatttgt ttaacctcat ttataaaagc ttcaaaaaaa 3550

ccca 3554

<210> 20

<211> 310

<212> PRT

<213> Homo sapiens

<400> 20

Met Ala Leu Arg Arg Pro Pro Arg Leu Arg Leu Cys Ala Arg Leu Pro Asp Phe Phe Leu Leu Leu Phe Arg Gly Cys Leu Ile Gly Ala Val Asn Leu Lys Ser Ser Asn Arg Thr Pro Val Val Gln Glu Phe Glu Ser Val Glu Leu Ser Cys Ile Ile Thr Asp Ser Gln Thr Ser Asp Pro Arg Ile Glu Trp Lys Lys Ile Gln Asp Glu Gln Thr Thr Tyr Val Phe Phe Asp Asn Lys Ile Gln Gly Asp Leu Ala Gly Arg Ala Glu Ile Leu Gly Lys Thr Ser Leu Lys Ile Trp Asn Val Thr Arg Arg Asp Ser Ala Leu Tyr Arg Cys Glu Val Val Ala Arg Asn Asp Arg Lys Glu Ile Asp Glu Ile Val Ile Glu Leu Thr Val 125 Gln Val Lys Pro Val Thr Pro Val Cys Arg Val Pro Lys Ala Val Pro Val Gly Lys Met Ala Thr Leu His Cys Gln Glu Ser Glu Gly His Pro Arg Pro His Tyr Ser Trp Tyr Arg Asn Asp Val Pro Leu Pro Thr Asp Ser Arg Ala Asn Pro Arg Phe Arg Asn Ser Ser Phe His Leu Asn Ser Glu Thr Gly Thr Leu Val Phe Thr Ala Val His Lys Asp Asp Ser Gly Gln Tyr Tyr Cys Ile Ala Ser Asn Asp Ala Gly Ser Ala Arg Cys Glu Glu Glu Met Glu Val Tyr Asp Leu Asn Ile Gly Gly Ile Ile Gly Gly Val Leu Val Val Leu Ala Val Leu Ala Leu Ile Thr Leu Gly Ile Cys Cys Ala Tyr Arg Arg Gly Tyr Phe Ile Asn Asn Lys Gln Asp Gly Glu Ser Tyr Lys Asn Pro Gly Lys Pro Asp Gly Val Asn Tyr Ile Arg Thr Asp Glu Glu Gly

290 295 300

Asp Phe Arg His Lys Ser Ser Phe Val Ile 305

<210> 21

<211> 3437

<212> DNA

<213> Homo sapiens

<400> 21

caggaccagg tettectacg etggageage ggggagaeag ecaecatgea 50 catcetegtg gtecatgeca tggtgatect getgacgetg ggeeegeete 100

gageegaega cagegagtte caggegetge tggacatetg gttteeggag 150 gagaagecae tgcccaccge etteetggtg gacacatcgg aggaggeget 200

gctgcttcct gactggctga agctgcgcat gatccgttct gaggtgctcc 250

geotggtgga egeegeeetg caggacetgg ageegeagea getgetgetg 300

ttegtgeagt egtttggeat eeeegtgtee ageatgagea aacteeteea 350

gttcctggac caggcagtgg cccacgaccc ccagactctg gagcagaaca 400

tcatggacaa gaattacatg geccaectgg tggaggteca geatgagege 450

ggcgcctccg gaggccagac tttccactcc ttgctcacag cctccctgcc 500

geccegeega gacageacag aggeaceeaa accaaagage ageeeagage 550

ageceatagg ceagggeegg attegggtgg ggaeceaget cegggtgetg 600

ggccctgagg acgacctggc tggcatgttc ctccagattt tcccgctcag 650 ecoggaccot eggtggeaga getecagtec ecgeceegtg gecetegeee 700

tgcagcagge cetgggeeag gagetggeee gegtegteea gggeageeee 750

gaggtgeegg geateaeggt gegtgteetg eaggeeeteg ceaccetget 800

cageteecca caeggeggtg ceetggtgat gteeatgeac egtageeact 850

tectggeetg ecegetgetg egecagetet gecagtacea gegetgtgtg 900

ccacaggaca deggettete etegetette etgaaggtge teetgeagat 950

gctgcagtgg ctggacagcc ctggcgtgga gggcgggccc ctgcgggcac 1000

ageteaggat gettgeeage eaggeeteag eegggegeag geteagtgat 1050

gtgcgagggg ggctcctgcg cctggccgag gccctggcct tccgtcagga 1100

cctggaggtg gtcagctcca ccgtccgtgc cgtcatcgcc accctgaggt 1150

ctggggagca gtgcagcgtg gagccggacc tgatcagcaa agtcctccag 1200

gggctgatcg aggtgaggtc cccccacctg gaggagctgc tgactgcatt 1250 cttctctgcc actgcggatg ctgcctcccc gtttccagcc tgtaagcccg 1300 ttgtggtggt gageteeetg etgetgeagg aggaggagee eetggetggg 1350 gggaageegg gtgeggaegg tggeageetg gaggeegtge ggetggggee 1400 ctogtcaggo etcctagtgg actggctgga aatgctggac cccgaggtgg 1450 teageagetg eccegacetg eageteagge tgetettete eeggaggaag 1500 ggcaaaggte aggccaggt gccctcgtte cgtccctace tectgaccet 1550 cttcacgcat cagtccagct ggcccacact gcaccagtgc atccgagtcc 1600 tgctgggcaa gagcegggaa cagaggtteg acceptetge etetetggae 1650 ttoototggg cotgoatoca tgttootogo atotggoagg ggogggacoa 1700 gegeaceceg cagaagegge gggaggaget ggtgetgegg gtecagggee 1750 eggageteat cageetggtg gagetgatee tggeegagge ggagaegegg 1800 agocaggacg gggacacago cgcctgcago ctcatccagg cccggctgcc 1850 cctgctgctc agctgctgct gtggggacga tgagagtgtc aggaaggtga 1900 cggagcacct gtcaggctgc atccagcagt ggggagacag cgtgctggga 1950 aggegetgee gagacettet cetgeagete tacetacage ggeeggaget 2000 gegggtgeee gtgeetgagg teetaetgea eagegaaggg getgeeagea 2050 gcagcgtetg caagctggac ggaetcatec accgetteat caegeteett 2100 geggaeaeca gegaeteeeg ggegttggag aacegagggg eggatgeeag 2150 catggeetge eggaagetgg eggtggegea eeegetgetg etgeteagge 2200 acctgcccat gategeggeg etectgeacg geegeaceca ceteaactte 2250 caggagttee ggeageagaa ceaeetgage tgetteetge aegtgetggg 2300 cctgctggag ctgctgcagc cgcacgtgtt ccgcagcgag caccaggggg 2350 cgctgtggga ctgccttctg tccttcatcc gcctgctgct gaattacagg 2400 aagtecteee gecatetgge tgeetteate aacaagtttg tgeagtteat 2450 ccataagtac attacctaca atgecceage agecatetee tteetgeaga 2500 agcacgccga cccgctccac gacctgtcct tcgacaacag tgacctggtg 2550 atgetgaaat eesteettge agggetsags stgessagsa gggasgasag 2600 gaccgaccga ggcctggacg aagagggcga ggaggagagc tcagccggct 2650 cettqccct qqtcaqcqte tccctgttca ccctctgac cgcggccgag 2700 atggcccct acatgaaacg gctttcccgg ggccaaacgg tggaggatct 2750 getggaggtt etgagtgaca tagaegagat gteeeggegg agaeeegaga 2800 tectgagett ettetegace aacetgeage ggetgatgag eteggeegag 2850 gaqtqttgcc gcaacctcgc cttcagcctg gccctgcgct ccatgcagaa 2900 cagecceage attgeagecg ettteetgee caegtteatg tactgeetgg 2950 geaqceagga ctttgaggtg gtgcagacgg ccctccggaa cctgcctgag 3000 tacqctctcc tgtgccaaga gcacgcggct gtgctgctcc accgggcctt 3050 cetggtggge atgtacggce agatggacce cagegegeag ateteegagg 3100 ccctqaqqat cctgcatatg gaggccgtga tgtgagcctg tggcagccga 3150 cocceteca ageocoggee egtecegtee coggggatee tegaggeaaa 3200 geccaggaag egtgggegtt getggtetgt eegaggaggt gagggegeeg 3250 agecetgagg ccaggeagge ccaggageaa tacteegage cctggggtgg 3300 ctccqggccg gccgctggca tcaggggccg tccagcaagc cctcattcac 3350 ettetgggee acageeetge egeggagegg eggateeece egggeatgge 3400 ctgggctggt tttgaatgaa acgacctgaa ctgtcaa 3437

<210> 22

<211> 1029

<212> PRT

<213> Homo sapiens

<400> 22

Met His Ile Leu Val Val His Ala Met Val Ile Leu Leu Thr Leu 1 5 10

Gly Pro Pro Arg Ala Asp Asp Ser Glu Phe Gln Ala Leu Leu Asp 20 25 30

Ile Trp Phe Pro Glu Glu Lys Pro Leu Pro Thr Ala Phe Leu Val 35 40 45

Asp Thr Ser Glu Glu Ala Leu Leu Leu Pro Asp Trp Leu Lys Leu
50
60

Arg Met Ile Arg Ser Glu Val Leu Arg Leu Val Asp Ala Ala Leu 65 70 75

Gln Asp Leu Glu Pro Gln Gln Leu Leu Phe Val Gln Ser Phe 80 85 90

Gly Ile Pro Val Ser Ser Met Ser Lys Leu Gln Phe Leu Asp 95 100 105

Gln	Ala	Val	Ala	His 110	Asp	Pro	Gln	Thr	Leu 115	Glu	Gln	Asn	Ile	Met 120
Asp	Lys	Asn	Tyr	Met 125	Ala	His	Leu	Val	Glu 130	Val	Gln	His	Glu	Arg 135
Gly	Ala	Ser	Gly	Gly 140	Gln	Thr	Phe	His	Ser 145	Leu	Leu	Thr	Ala	Ser 150
Leu	Pro	Pro	Arg	Arg 155	Asp	Ser	Thr	Glu	Ala 160	Pro	Lys	Pro	Lys	Ser 165
Ser	Pro	Glu	Gln	Pro 170	Ile	Gly	Gln	Gly	Arg 175	Ile	Arg	Val	Gly	Thr 180
Gln	Leu	Arg	Val	Leu 185	Gly	Pro	Glu	Asp	Asp 190	Leu	Ala	Gly	Met	Phe 195
Leu	Gln	Ile	Phe	Pro 200	Leu	Ser	Pro	Asp	Pro 205	Arg	Trp	Gln	Ser	Ser 210
Ser	Pro	Arg	Pro	Val 215	Ala	Leu	Ala	Leu	Gln 220	Gln	Ala	Leu	Gly	Gln 225
Glu	Leu	Ala	Arg	Val 230	Val	Gln	Gly	Ser	Pro 235	Glu	Val	Pro	Gly	Ile 240
Thr	Val	Arg	Val	Leu 245	Gln	Ala	Leu	Ala	Thr 250	Leu	Leu	Ser	Ser	Pro 255
His	Gly	Gly	Ala	Leu 260	Val	Met	Ser	Met	His 265	Arg	Ser	His	Phe	Leu 270
Ala	Сув	Pro	Leu	Leu 275	Arg	Gln	Leu	Cys	Gln 280	Tyr	Gln	Arg	Cys	Val 285
Pro	Gln	Asp	Thr	Gly 290	Phe	Ser	Ser	Leu	Phe 295	Leu	Lys	Val	Leu	Leu 300
Gln	Met	Leu	Gln	Trp 305	Leu	Asp	Ser	Pro	Gly 310	Val	Glu	Gly	Gly	Pro 315
Leu	Arg	Ala	Gln	Leu 320	Arg	Met	Leu	Ala	Ser 325	Gln	Ala	Ser	Ala	Gly 330
Arg	Arg	Leu	Ser	Asp 335	Val	Arg	Gly	Gly	Leu 340	Leu	Arg	Leu	Ala	Glu 345
Ala	Leu	Ala	Phe	Arg 350	Gln	Asp	Leu	Glu	Val 355	Val	Ser	Ser	Thr	Val 360
Arg	Ala	Val	Ile	Ala 365	Thr	Leu	Arg	Ser	Gly 370	Glu	Gln	Cys	Ser	Val 375
Glu	Pro	Asp	Leu	Ile 380	Ser	Lys	Val	Leu	Gln 385	Gly	Leu	Ile	Glu	Val 390
Arg	Ser	Pro	His	Leu	Glu	Glu	Leu	Leu	Thr	Ala	Phe	Phe	Ser	Ala

				395					400					405
Thr	Ala	Asp	Ala	Ala 410	Ser	Pro	Phe	Pro	Ala 415	Cys	Lys	Pro	Val	Val 420
Val	Val	Ser	Ser	Leu 425	Leu	Leu	Gln	Glu	Glu 430	Glu	Pro	Leu	Ala	Gly 435
Gly	Lys	Pro	Gly	Ala 440	Asp	Gly	Gly	Ser	Leu 445	Glu	Ala	Val	Arg	Leu 450
Gly	Pro	Ser	Ser	Gly 455	Leu	Leu	Val	Asp	Trp 460	Leu	Glu	Met	Leu	Asp 465
Pro	Glu	Val	Val	Ser 470	Ser	Cys	Pro	Asp	Leu 475	Gln	Leu	Arg	Leu	Leu 480
Phe	Ser	Arg	Arg	Lys 485	Gly	Lys	Gly	Gln	Ala 490	Gln	Val	Pro	Ser	Phe 495
Arg	Pro	Tyr	Leu	Leu 500	Thr	Leu	Phe	Thr	His 505	Gln	Ser	Ser	Trp	Pro 510
Thr	Leu	His	Gln	Cys 515	Ile	Arg	Val	Leu	Leu 520	Gly	Lys	Ser	Arg	Glu 525
Gln	Arg	Phe	Asp	Pro 530	Ser	Ala	Ser	Leu	Asp 535	Phe	Leu	Trp	Ala	Cys 540
Ile	His	Val	Pro	Arg 545	Ile	Trp	Gln	Gly	Arg 550	Asp	Gln	Arg	Thr	Pro 555
Gln	Lys	Arg	Arg	Glu 560	Glu	Leu	Val	Leu	Arg 565	Val	Gln	Gly	Pro	Glu 570
Leu	Ile	Ser	Leu	Val 575	Glu	Leu	Ile	Leu	Ala 580	Glu	Ala	Glu	Thr	Arg 585
Ser	Gln	Asp	Gly	Asp 590	Thr	Ala	Ala	Cys	Ser 595	Leu	Ile	Gln	Ala	Arg 600
Leu	Pro	Leu	Leu	Leu 605	Ser	Cys	Cys	Cys	Gly 610	Asp	Asp	Glu	Ser	Val 615
Arg	Lys	Val	Thr	Glu 620	His	Leu	Ser	Gly	Cys 625	Ile	Gln	Gln	Trp	Gly 630
Asp	Ser	Val	Leu	Gly 635	Arg	Arg	Cys	Arg	Asp 640	Leu	Leu	Leu	Gln	Leu 645
Tyr	Leu	Gln	Arg	Pro 650	Glu	Leu	Arg	Val	Pro 655	Val	Pro	Glu	Val	Leu 660
Leu	His	Ser	Glu	Gly 665	Ala	Ala	Ser	Ser	Ser 670	Val	Cys	Lys	Leu	Asp 675
Gly	Leu	Ile	His	Arg 680	Phe	Ile	Thr	Leu	Leu 685	Ala	Asp	Thr	Ser	Asp 690

Ser Arg Ala Leu Glu Asn Arg Gly Ala Asp Ala Ser Met Ala Cys Arg Lys Leu Ala Val Ala His Pro Leu Leu Leu Arg His Leu Pro Met Ile Ala Ala Leu Leu His Gly Arg Thr His Leu Asn Phe Gln Glu Phe Arg Gln Gln Asn His Leu Ser Cys Phe Leu His Val 745 750 Leu Gly Leu Leu Glu Leu Leu Gln Pro His Val Phe Arg Ser Glu His Gln Gly Ala Leu Trp Asp Cys Leu Leu Ser Phe Ile Arg Leu Leu Leu Asn Tyr Arg Lys Ser Ser Arg His Leu Ala Ala Phe Ile Asn Lys Phe Val Gln Phe Ile His Lys Tyr Ile Thr Tyr Asn Ala Pro Ala Ala Ile Ser Phe Leu Gln Lys His Ala Asp Pro Leu His 815 Asp Leu Ser Phe Asp Asn Ser Asp Leu Val Met Leu Lys Ser Leu Leu Ala Gly Leu Ser Leu Pro Ser Arg Asp Asp Arg Thr Asp Arg 845 Gly Leu Asp Glu Glu Glu Glu Glu Ser Ser Ala Gly Ser Leu Pro Leu Val Ser Val Ser Leu Phe Thr Pro Leu Thr Ala Ala Glu Met Ala Pro Tyr Met Lys Arg Leu Ser Arg Gly Gln Thr Val Glu Asp Leu Leu Glu Val Leu Ser Asp Ile Asp Glu Met Ser Arg Arg Arg Pro Glu Ile Leu Ser Phe Phe Ser Thr Asn Leu Gln Arg Leu 920 925 Met Ser Ser Ala Glu Glu Cys Cys Arg Asn Leu Ala Phe Ser Leu 940 Ala Leu Arg Ser Met Gln Asn Ser Pro Ser Ile Ala Ala Ala Phe Leu Pro Thr Phe Met Tyr Cys Leu Gly Ser Gln Asp Phe Glu Val Val Gln Thr Ala Leu Arg Asn Leu Pro Glu Tyr Ala Leu Leu Cys 980 985 990

Gln Glu His Ala Ala Val Leu Leu His Arg Ala Phe Leu Val Gly
995 1000 1005

Met Tyr Gly Gln Met Asp Pro Ser Ala Gln Ile Ser Glu Ala Leu 1010 1015 1020

Arg Ile Leu His Met Glu Ala Val Met 1025

<210> 23

<211> 2186

<212> DNA

<213> Homo sapiens

<400> 23

cegggecatg cageetegge eeegegggeg eeegeegege aeeegaggag 50 atgaggetee geaatggeac etteetgaeg etgetgetet tetgeetgtg 100 cgecttecte tegetgteet ggtacgegge acteagegge cagaaaggeg 150 acgttgtgga cgtttaccag cgggagttcc tggcgctgcg cgatcggttg 200 cacgcagetg ageaggagag ceteaagege tecaaggage teaacetggt 250 getggaegag ateaagaggg eegtgteaga aaggeaggeg etgegagaeg 300 gagacggcaa tcgcacctgg ggccgcctaa cagaggaccc ccgattgaag 350 cogtggaacg getcacaccg geacgtgetg cacetgeeca cogtetteea 400 tcacctgcca cacctgctgg ccaaggagag cagtctgcag cccgcggtgc 450 gegtgggcca gggccgcacc ggagtgtcgg tggtgatggg catcccgagc 500 qtqcqgcgcg aggtgcactc gtacctgact gacactctgc actcgctcat 550 ctccgagctg agcccgcagg agaaggagga ctcggtcatc gtggtgctga 600 tegeogagae tgaeteaeag tacaettegg cagtgacaga gaacatcaag 650 gccttgttcc ccacggagat ccattctggg ctcctggagg tcatctcacc 700 ctcccccae ttctaccctg acttctcccg cctccgagag tcctttgggg 750 accecaagga gagagteagg tggaggaeea aacagaacet egattactge 800 ttcctcatga tgtacgcgca gtccaaaggc atctactacg tgcagctgga 850 ggatgacate gtggccaage ccaactacet gagcaccatg aagaactttg 900 cactgoagoa qoottoagag gactggatga tootqgagtt otoccagotg 950 ggetteattg gtaagatgtt caagtegetg gaeetgagee tgattgtaga 1000 gttcattctc atgttctacc gggacaagcc catcgactgg ctcctggacc 1050

atattetqtq qqtqaaaqte tqcaaceceq aqaaggatge gaaqcactgt 1100 gaccggcaga aagccaacct gcggatccgc ttcaaaccgt ccctcttcca 1150 gcacgtgggc actcactcct cgctggctgg caagatccag aaactgaagg 1200 acaaaqactt tqqaaaqcaq qcqctqcqqa agqaqcatqt gaacccqcca 1250 quaqaqqtqa quaqqqct gaagacatac cagcacttca coctggagaa 1300 agoctacetg egegaggaet tettetggge etteacecet geegeggggg 1350 actteateeg etteegette tteeaaeete taagaetgga geggttette 1400 ttccqcaqtq qqaacatcga gcacccggag gacaagctct tcaacacgtc 1450 tgtggaggtg ctgcccttcg acaaccctca gtcagacaag gaggccctgc 1500 aggagggeeg cacegeeace eteeggtace eteggageee egaeggetae 1550 ctccagateg geteetteta caagggagtg geagagggag aggtggaeec 1600 agoettegge cetetggaag caetgegeet etegateeag aeggaeteee 1650 etgtgtgggt gattetgage gagatettee tgaaaaagge egaetaaget 1700 qeqqqettet qaqqqtaeec tgtgqeeage cetgaageec acatttetgg 1750 gggtgtcgtc actgccgtcc ccggagggcc agatacggcc ccgcccaaag 1800 ggttctgcct ggcgtcgggc ttgggccggc ctggggtccg ccgctggccc 1850 ggaggcccta ggagctggtg ctgccccgc ccgccgggcc gcggaggagg 1900 caggeggeec ceacactgtg cetgaggeec ggaaccgtte geacceggee 1950 tgccccagtc aggccgtttt agaagagctt ttacttgggc gcccgccgtc 2000 tetggegega acaetggaat geatataeta etttatgtge tgtgtttttt 2050 attettggat acatttgatt ttttcacgta agtecacata tacttetata 2100 aaaaaaaaaa aaaaaaaaaa aaaaaaaaa aaaaaa 2186

<210> 24

<211> 548

<212> PRT

<213> Homo sapiens

<400> 24

Met Arg Leu Arg Asn Gly Thr Phe Leu Thr Leu Leu Leu Phe Cys 1 5 10 15

Leu Cys Ala Phe Leu Ser Leu Ser Trp Tyr Ala Ala Leu Ser Gly
20 25 30

Gln Lys Gly Asp Val Val Asp Val Tyr Gln Arg Glu Phe Leu Ala Leu Arg Asp Arg Leu His Ala Ala Glu Gln Glu Ser Leu Lys Arg Ser Lys Glu Leu Asn Leu Val Leu Asp Glu Ile Lys Arg Ala Val Ser Glu Arg Gln Ala Leu Arg Asp Gly Asp Gly Asn Arg Thr Trp Gly Arg Leu Thr Glu Asp Pro Arg Leu Lys Pro Trp Asn Gly Ser His Arg His Val Leu His Leu Pro Thr Val Phe His His Leu Pro 110 His Leu Leu Ala Lys Glu Ser Ser Leu Gln Pro Ala Val Arg Val 125 130 Gly Gln Gly Arg Thr Gly Val Ser Val Val Met Gly Ile Pro Ser Val Arg Arg Glu Val His Ser Tyr Leu Thr Asp Thr Leu His Ser Leu Ile Ser Glu Leu Ser Pro Gln Glu Lys Glu Asp Ser Val Ile 170 Val Val Leu Ile Ala Glu Thr Asp Ser Gln Tyr Thr Ser Ala Val Thr Glu Asn Ile Lys Ala Leu Phe Pro Thr Glu Ile His Ser Gly Leu Leu Glu Val Ile Ser Pro Ser Pro His Phe Tyr Pro Asp Phe 215 220 Ser Arg Leu Arg Glu Ser Phe Gly Asp Pro Lys Glu Arg Val Arg Trp Arg Thr Lys Gln Asn Leu Asp Tyr Cys Phe Leu Met Met Tyr Ala Gln Ser Lys Gly Ile Tyr Tyr Val Gln Leu Glu Asp Asp Ile 265 Val Ala Lys Pro Asn Tyr Leu Ser Thr Met Lys Asn Phe Ala Leu Gln Gln Pro Ser Glu Asp Trp Met Ile Leu Glu Phe Ser Gln Leu Gly Phe Ile Gly Lys Met Phe Lys Ser Leu Asp Leu Ser Leu Ile 310 Val Glu Phe Ile Leu Met Phe Tyr Arg Asp Lys Pro Ile Asp Trp

				320					325					330
Leu	Leu	Asp	His	Ile 335	Leu	Trp	Val	Lys	Val 340	Cys	Asn	Pro	Glu	Lys 345
Asp	Ala	Lys	His	Сув 350	Asp	Arg	Gln	Lys	Ala 355	Asn	Leu	Arg	Ile	Arg 360
Phe	Lys	Pro	Ser	Leu 365	Phe	Gln	His	Val	Gly 370	Thr	His	Ser	Ser	Leu 375
Ala	Gly	Lys	Ile	Gln 380	Lys	Leu	Lys	Asp	Lys 385	Asp	Phe	Gly	Lys	Gln 390
Ala	Leu	Arg	Lys	Glu 395	His	Val	Asn	Pro	Pro 400	Ala	Glu	Val	Ser	Thr 405
Ser	Leu	Lys	Thr	Tyr 410	Gln	His	Phe	Thr	Leu 415	Glu	Lys	Ala	Tyr	Leu 420
Arg	Glu	Asp	Phe	Phe 425	Trp	Ala	Phe	Thr	Pro 430	Ala	Ala	Gly	Asp	Phe 435
Ile	Arg	Phe	Arg	Phe 440	Phe	Gln	Pro	Leu	Arg 445	Leu	Glu	Arg	Phe	Phe 450
Phe	Arg	Ser	Gly	Asn 455	Ile	Glu	His	Pro	Glu 460	Asp	Lys	Leu	Phe	Asn 465
Thr	Ser	Val	Glu	Val 470	Leu	Pro	Phe	Asp	Asn 475	Pro	Gln	Ser	Asp	Lys 480
Glu	Ala	Leu	Gln	Glu 485	Gly	Arg	Thr	Ala	Thr 490	Leu	Arg	Tyr	Pro	Arg 495
Ser	Pro	Asp	Gly	Tyr 500	Leu	Gln	Ile	Gly	Ser 505	Phe	Tyr	Lys	Gly	Val 510
Ala	Glu	Gly	Glu	Val 515	Asp	Pro	Ala	Phe	Gly 520	Pro	Leu	Glu	Ala	Leu 525
Arg	Leu	Ser	Ile	Gln 530	Thr	Asp	Ser	Pro	Val 535	Trp	Val	Ile	Leu	Ser 540
Glu	Ile	Phe	Leu	Lys 545	Lys	Ala	Asp							
<210>	- 25													
<211>														
<212> <213>			cial	Sequ	ience	3								
				-										
<220> <223>		thet	ic C	ligo	onucl	eoti	.de E	robe	è					
<400>	25													
tgta	aaac	ga c	ggcc	agtt	a aa	taga	ccto	caa	ittat	taa	tct	43		

```
The flash film draw great the film of the flash the flash than the
```

```
<210> 26
<211> 41
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
 caggaaacag ctatgaccac ctgcacacct gcaaatccat t 41
<210> 27
<211> 19
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 27
actogggatt cotgetgtt 19
<210> 28
<211> 23
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 28
aggeetttae eeaaggeeae aac 23
<210> 29
<211> 19
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 29
ggcctgtcct gtgttctca 19
<210> 30
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 30
tcccaccact tacttccatg aa 22
<210> 31
<211> 25
<212> DNA
```

```
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 31
 ctgtggtacc caattgccgc cttgt 25
<210> 32
<211> 23
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 32
 attgtcctga gattcgagca aga 23
<210> 33
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 33
 gtccagcaag ccctcatt 18
<210> 34
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 34
 cttctgggcc acagccctgc 20
<210> 35
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 35
cagttcaggt cgtttcattc a 21
<210> 36
<211> 19
<212> DNA
<213> Artificial Sequence
<220>
```



- <223> Synthetic Oligonucleotide Probe
- <400> 36 ccagtcaggc cgttttaga 19
- <210> 37
- <211> 21
- <212> DNA
- <213> Artificial Sequence
- <220s
- <223> Synthetic Oligonucleotide Probe
- <400> 37
- cgggcgccca agtaaaagct c 21
- <210> 38
- <211> 28
- <212> DNA
- <213> Artificial Sequence
- <2200 S
- <223> Synthetic Oligonucleotide Probe
- <400> 38
- cataaagtag tatatgcatt ccagtgtt 28